> This Raw Listing contains the General Information
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 1   <110> APPLICANT: Kawasaki, Hiroaki
 2         Graybiel, Ann
 3         Housman, David
 4   <120> TITLE OF INVENTION: Genes Integrating Signal Transduction Pathways
 5   <130> FILE REFERENCE: MIT-103
 6   <140> CURRENT APPLICATION NUMBER: US/09/422,999
 7   <141> CURRENT FILING DATE: 1999-10-22
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11   <151> EARLIER FILING DATE: 1998-11-16
12   <160> NUMBER OF SEQ ID NOS: 18
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35       Arg Met Phe Leu Met Met His Pro Trp Tyr Ile Pro Ser Ser Gln Leu
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37       gct tcg aaa ctg ctc cac ttc tat cag caa tcc cgg aag gac aac tcc    369
38       Ala Ser Lys Leu Leu His Phe Tyr Gln Gln Ser Arg Lys Asp Asn Ser
39                   55                  60                  65
40       aat tcc cta cag gtg aaa acc tgt cac ctg gtc agg tac tgg gtc tca    417
41       Asn Ser Leu Gln Val Lys Thr Cys His Leu Val Arg Tyr Trp Val Ser
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47    Lys Glu Leu Lys Ala Leu Leu Asp Gln Glu Gly Asn Arg Arg His Ser
48                    105                 110                 115
49    agc ctc atc gac atc gag agt gtc ccc acc tac aag tgg aag cgg cag   561
50    Ser Leu Ile Asp Ile Glu Ser Val Pro Thr Tyr Lys Trp Lys Arg Gln
51                    120                 125                 130
52    gtg acc cag cgg aac cct gtg gaa cag aaa aag cgc aag atg tcc ctg   609
53    Val Thr Gln Arg Asn Pro Val Glu Gln Lys Lys Arg Lys Met Ser Leu
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55    ttg ttt gat cac ttg gag cct atg gaa ctg gca gaa cat ctc acc tac   657
56    Leu Phe Asp His Leu Glu Pro Met Glu Leu Ala Glu His Leu Thr Tyr
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59    Leu Glu Tyr Arg Ser Phe Cys Lys Ile Leu Phe Gln Asp Tyr His Ser
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61    ttt gtg act cat ggc tgc act gta gac aat ccg gtc ctg gag cga ttc   753
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68    Ser Lys Pro Thr Ala Thr Gln Arg Ala Leu Val Ile Thr His Phe Val
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70    cat gtg gca gag aag ctg ctg cag ctg cag aac ttc aac acg ttg atg   897
71    His Val Ala Glu Lys Leu Leu Gln Leu Gln Asn Phe Asn Thr Leu Met
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73    gcc gtc gtg gga ggc ctg agc cac agc tcc atc tca cgc ctc aag gag   945
74    Ala Val Val Gly Gly Leu Ser His Ser Ser Ile Ser Arg Leu Lys Glu
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76    acc cac agc cat gtc agc cct gac acc atc aag ctc tgg gaa ggt ctg   993
77    Thr His Ser His Val Ser Pro Asp Thr Ile Lys Leu Trp Glu Gly Leu
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79    aca gaa cta gtg aca gct act ggc aac tac agc aac tac cgg cga agg   1041
80    Thr Glu Leu Val Thr Ala Thr Gly Asn Tyr Ser Asn Tyr Arg Arg Arg
81                    280                 285                 290
82    ctg gcg gcc tgc gtg ggc ttc cgc ttt cct atc ctg ggt gtg cac ctc   1089
83    Leu Ala Ala Cys Val Gly Phe Arg Phe Pro Ile Leu Gly Val His Leu
84                    295                 300                 305
85    aag gat cta gtg gct ctg cag ctg gct ctg cct gac tgg ctg gac cca   1137
86    Lys Asp Leu Val Ala Leu Gln Leu Ala Leu Pro Asp Trp Leu Asp Pro
87              310                 315                 320
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89    Gly Arg Thr Arg Leu Asn Gly Ala Lys Met Arg Gln Leu Phe Ser Ile
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91    ctg gag gag ttg gct atg gtg acc agt ctt cga cca cca gtg caa gcc   1233
92    Leu Glu Glu Leu Ala Met Val Thr Ser Leu Arg Pro Pro Val Gln Ala
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100        aag tca tcg ccc acc agc ccc acc agc tgc acc ccg cct ccc cgg ccg    1377
101        Lys Ser Ser Pro Thr Ser Pro Thr Ser Cys Thr Pro Pro Pro Arg Pro
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104        Pro Val Leu Glu Glu Trp Thr Ser Val Ala Lys Pro Lys Leu Asp Gln
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109        aac ttt gac gtt gat ggg gac ggt cac atc tcc cag gag gag ttc cag    1521
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114                      455                 460                 465
115        cag aac cag gat ggc tgc atc agc cgg gag gag atg att tcc tac ttc    1617
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143        ggattctggg gcaggaggtt ggggctgaag gtgggggaag ttgaaggtgg catgcactga  2119
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172      His Leu Thr Tyr Leu Glu Tyr Arg Ser Phe Cys Lys Ile Leu Phe Gln
173                          165                 170                 175
174      Asp Tyr His Ser Phe Val Thr His Gly Cys Thr Val Asp Asn Pro Val
175                      180                 185                 190
176      Leu Glu Arg Phe Ile Ser Leu Phe Asn Ser Val Ser Gln Trp Val Gln
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178      Leu Met Ile Leu Ser Lys Pro Thr Ala Thr Gln Arg Ala Leu Val Ile
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184      Arg Leu Lys Glu Thr His Ser His Val Ser Pro Asp Thr Ile Lys Leu
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186      Trp Glu Gly Leu Thr Glu Leu Val Thr Ala Thr Gly Asn Tyr Ser Asn
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190      Gly Val His Leu Lys Asp Leu Val Ala Leu Gln Leu Ala Leu Pro Asp
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199                    370                 375                 380
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202      Pro Pro Arg Pro Pro Val Leu Glu Glu Trp Thr Ser Val Ala Lys Pro
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204      Lys Leu Asp Gln Ala Leu Val Ala Glu His Ile Glu Lys Met Val Glu
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206      Ser Val Phe Arg Asn Phe Asp Val Asp Gly Asp Gly His Ile Ser Gln
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208      Glu Glu Phe Gln Ile Ile Arg Gly Asn Phe Pro Tyr Leu Ser Ala Phe
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